RAW SEQUENCE LISTING   DATE: 04/23/2002
PATENT APPLICATION: US/09/845,612B   TIME: 08:16:31

Input Set : A:\UTSD795.txt
Output Set: N:\CRF3\04232002\I845612B.raw

     3 <110> APPLICANT: YU, HONGTAO
     4   TANG, ZHANYUN
     5   LUO, XUELIAN
     6   RIZO-REY, JOSE
     8 <120> TITLE OF INVENTION: A CLASS OF 12MER PEPTIDES THAT INHIBIT THE FUNCTINO OF THE
MITOTIC CHECK
     9   POINT PROTEIN MAD2
    11 <130> FILE REFERENCE: UTSD:795
    13 <140> CURRENT APPLICATION NUMBER: 09/845,612B
    14 <141> CURRENT FILING DATE: 2001-04-27
    16 <160> NUMBER OF SEQ ID NOS: 20
    18 <170> SOFTWARE: PatentIn version 3.0

ERRORED SEQUENCES

   333 <210> SEQ ID NO: 20
   334 <211> LENGTH: 11
   335 <212> TYPE: PRT
   336 <213> ORGANISM: HIV-TAT PROTEIN
   338 <400> SEQUENCE: 20
   340 Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg
   341 1   5   10
   343 25025713.1
E--> 344 4
E--> 345 Express Mail No. EH244916502US   — delete
E--> 348 7
E--> 356 UTSD795.txt

4/23/02

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 04/23/2002
PATENT APPLICATION: US/09/845,612B   TIME: 08:16:33

Input Set : A:\UTSD795.txt
Output Set: N:\CRF3\04232002\I845612B.raw

Invalid <213> Response:
Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1

VERIFICATION SUMMARY   DATE: 04/23/2002
PATENT APPLICATION: US/09/845,612B   TIME: 08:16:33

          Input Set : A:\UTSD795.txt
          Output Set: N:\CRF3\04232002\I845612B.raw

L:23 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
L:42 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:292 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
L:314 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:344 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:20
L:345 M:333 E: Wrong sequence grouping, Amino acids not in groups!
L:345 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:4
M:332 Repeated in SeqNo=20
L:356 M:333 E: Wrong sequence grouping, Amino acids not in groups!
L:356 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:1
L:356 M:252 E: No. of Seq. differs, <211> LENGTH:Input:11 Found:16 SEQ:20